## 2) 2002 E. Steven L. Oh, Esther H. Walsh, Michael

## SEQUENCE LISTING

<120> Methods of Diagnosing Liver Fibrosis

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<151> 2002-02-28

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atc tgt gca aac ggg cgg caa act gtg tcc tgg gca gta acc cca aag 96
Ile Cys Ala Asn Gly Arg Gln Thr Val Ser Trp Ala Val Thr Pro Lys
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Ser Leu Gly Asn Val Asn Phe Thr Val Ser Ala Glu Ala Leu Glu Ser
35 40 45

caa gag ctg tgt ggg act gag gtg cct tca gtt cct gaa cac gga agg 192 Gln Glu Leu Cys Gly Thr Glu Val Pro Ser Val Pro Glu His Gly Arg 50 55 60

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Lys Asp Thr Val Ile Lys Pro Leu Leu Val Glu Pro Glu Gly Leu Glu
65 70 75 80

aag gaa aca aca ttc aac tcc cta ctt tgt cca tca ggt ggt gag gtt 288 Lys Glu Thr Thr Phe Asn Ser Leu Leu Cys Pro Ser Gly Gly Glu Val

90

RECEIVED TO MAIL ROW.

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Ala	Arg	Ala 115	Ser	Val	Ser	Val	Leu 120	Gly	Asp	Ile	Leu	Gly 125	Ser	Ala	Met	
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	130					135					140					
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Asn 145	мет	vaı	Leu	Pne	150	Pro	ASII	ше	Tyr	155	Leu	Asp	ıyı	цец	160	
gaa	aca	caq	caq	ctt	act	сса	gag	atc	aag	tcc	aag	gcc	att	ggc	tat	528
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Ser	Tyr	195	Thr	Pne	GIÀ	GIU	200	Tyr	Gly	Arg	ASII	205	GIÀ	ASII	1111	
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									Phe							
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															tcc Ser	720
225					230					235					240	
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GIII	Arg	GIII	ьуѕ	245	ASII	GIY	Cys	PHE	250	261	261	Gry	361	255		
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		Thr					Glu					Val	Thr		ect Pro	004
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val	290	_	noil	ATA	ъси	295	Cys	neu	Ų1u	Der	300		<b>-</b> 75			
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acc Thr	aac Asn	atc Ile	gtg Val	aag Lys 405	tgg Trp	atc Ile	acg Thr	aag Lys	cag Gln 410	cag Gln	aat Asn	gcc Ala	cag Gln	ggc Gly 415	ggt Gly	1248
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tat Tyr	gga Gly	gca Ala 435	gcc Ala	aca Thr	ttt Phe	acc Thr	agg Arg 440	act Thr	Gly	aag Lys	gct Ala	gca Ala 445	cag Gln	gtg Val	act Thr	1344
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ttg Leu	aaa Lys	tac Tyr	aat Asn 500	att Ile	ctc Leu	cca Pro	gaa Glu	aag Lys 505	Glu	gag Glu	ttc Phe	ccc Pro	ttt Phe 510	Ala	tta Leu	1536
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		_	cct tgc agc aaa gat ct Pro Cys Ser Lys Asp Let 635 64	u
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Gln Glu Leu Cys 50	Gly Thr Glu 55	Val Pro Ser	Val Pro Glu His Gly Arg	3
Lys Asp Thr Val 65	Ile Lys Pro 70	Leu Leu Val	Glu Pro Glu Gly Leu Glu 75 80	1

Lys	Glu	Thi	Thi	Pho 85	e Asr	ı Ser	Leu	Lev	ι Cys 90	s Pro	o Ser	Gly	/ Gly	/ Gl: 95	ı Val
Ser	Glu	Glı	1 Let 100	ı Se:	r Let	Lys	Leu	Pro	Pro	Ası	ı Val	l Val	l Glu 110	ı Glı	ı Ser
Ala	Arg	Ala 115	a Ser	Va.	l Ser	· Val	. Leu 120	Gly	Asp	Ile	e Lev	Gly 125	/ Sei	Ala	a Met
Gln	Asn 130	Thi	Glr	Ası	ı Lev	Leu 135		Met	Pro	Туг	Gly	cys		/ Gli	Gln
Asn 145	Met	Val	. Leu	Phe	e Ala 150	Pro	Asn	Ile	Туг	7 Val	. Leu		Туг	Leu	Asn 160
				165	5				170	)				175	Tyr
			180					185					190	Asp	Gly
		195					200					205			Thr
	210					215					220				Tyr
225					230					235					Ser 240
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		275			Leu		280					285			
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305					Gly 310	•				315					320
				325					330					335	
			340		Glu			345					350		
		355			Pro		360					365			
	370					375					380				
Tyr 385	Leu	Thr	Ala	Gln	Pro 390	Ala	Pro	Thr	Ser	Glu 395	Asp	Leu	Thr	Ser	Ala 400
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			420		Asp			425					430		
Tyr		435					440					445			
	450				*	455					460				
Asn 2	Arg	Leu	Leu	Leu	Gln 470	Gln	Val	Ser	Leu	Pro 475	Glu	Leu	Pro	Gly	
Tyr	Ser 1	Met	Lys	Val 485		Gly	Glu		Cys 490		Tyr	Leu		Thr 495	480 Ser
Leu 1	Lys '	Tyr	Asn	Ile	Leu	Pro	Glu :			Glu	Phe	Pro	Phe	Ala	Leu

500 505 Gly Val Gln Thr Leu Pro Gln Thr Cys Asp Glu Pro Lys Ala His Thr 520 Ser Phe Gln Ile Ser Leu Ser Val Ser Tyr Thr Gly Ser Arg Ser Ala 535 540 Ser Asn Met Ala Ile Val Asp Val Lys Met Val Ser Gly Phe Ile Pro 550 555 Leu Lys Pro Thr Val Lys Met Leu Glu Arg Ser Asn His Val Ser Arg 565 570 Thr Glu Val Ser Ser Asn His Val Leu Ile Tyr Leu Asp Lys Val Ser 585 Asn Gln Thr Leu Ser Leu Phe Phe Thr Val Leu Gln Asp Val Pro Val 600 605 Arg Asp Leu Lys Pro Ala Ile Val Lys Val Tyr Asp Tyr Tyr Glu Thr 615 Asp Glu Phe Ala Ile Ala Glu Tyr Asn Ala Pro Cys Ser Lys Asp Leu 630 625 635 Gly Asn Ala <210> 3 <211> 782 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (63)...(686) <400> 3 aggggcctta gcgtgccgca tcgccgagat ccagcgccca gagagacacc agagaaccca 60 cc atg gcc ccc ttt gag ccc ctg gct tct ggc atc ctg ttg ttg ctg Met Ala Pro Phe Glu Pro Leu Ala Ser Gly Ile Leu Leu Leu 5 10 tgg ctg ata gcc ccc agc agg gcc tgc acc tgt gtc cca ccc cac cca 155 Trp Leu Ile Ala Pro Ser Arg Ala Cys Thr Cys Val Pro Pro His Pro 20 cag acg gcc ttc tgc aat tcc gac ctc gtc atc agg gcc aag ttc gtg 203 Gln Thr Ala Phe Cys Asn Ser Asp Leu Val Ile Arg Ala Lys Phe Val ggg aca cca gaa gtc aac cag acc acc tta tac cag cgt tat gag atc 251 Gly Thr Pro Glu Val Asn Gln Thr Thr Leu Tyr Gln Arg Tyr Glu Ile

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Lys Met Thr Lys Met Tyr Lys Gly Phe Gln Ala Leu Gly Asp Ala Ala

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75

70

65

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347

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		_	-										aag Lys			491
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_		_	_	_				_	_		_	_	cag Gln			587
			_	_			_		_			_	tgc Cys			635
				_				_					cag Gln 205			683
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tga * ctc <21 <21:		ott 1	tette	ccgga									gtto	cca		
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Ile Arg Phe Val Tyr Thr Pro Ala Met Glu Ser Val Cys Gly Tyr Phe His Arg Ser His Asn Arg Ser Glu Glu Phe Leu Ile Ala Gly Lys Leu Gln Asp Gly Leu Leu His Ile Thr Thr Cys Ser Phe Val Ala Pro Trp Asn Ser Leu Ser Leu Ala Gln Arg Arg Gly Phe Thr Lys Thr Tyr Thr Val Gly Cys Glu Glu Cys Thr Val Phe Pro Cys Leu Ser Ile Pro Cys Lys Leu Gln Ser Gly Thr His Cys Leu Trp Thr Asp Gln Leu Leu Gln Gly Ser Glu Lys Gly Phe Gln Ser Arg His Leu Ala Cys Leu Pro Arg Glu Pro Gly Leu Cys Thr Trp Gln Ser Leu Arg Ser Gln Ile Ala